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Sequence Listing was accepted.

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217-9197 (toll free).

Reviewer: Durreshwar Anjum

Timestamp: Mon Oct 15 09:17:02 EDT 2007

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Application No: 10560303 Version No: 1.0

Input Set:

Output Set:

Started: 2007-09-26 16:48:14.742
Finished: 2007-09-26 16:48:27.466
Elapsed: 0 hr(s) 0 min(s) 12 sec(s) 724 ms
Total Warnings: 60
Total Errors: 0
No. of SeqIDs Defined: 92
Actual SeqID Count: 92

Error code	Error Description
W 402	Undefined organism found in <213> in SEQ ID (1)
W 402	Undefined organism found in <213> in SEQ ID (2)
W 402	Undefined organism found in <213> in SEQ ID (3)
W 402	Undefined organism found in <213> in SEQ ID (4)
W 402	Undefined organism found in <213> in SEQ ID (5)
W 402	Undefined organism found in <213> in SEQ ID (6)
W 402	Undefined organism found in <213> in SEQ ID (7)
W 402	Undefined organism found in <213> in SEQ ID (8)
W 213	Artificial or Unknown found in <213> in SEQ ID (9)
W 213	Artificial or Unknown found in <213> in SEQ ID (10)
W 213	Artificial or Unknown found in <213> in SEQ ID (11)
W 213	Artificial or Unknown found in <213> in SEQ ID (12)
W 213	Artificial or Unknown found in <213> in SEQ ID (13)
W 213	Artificial or Unknown found in <213> in SEQ ID (14)
W 213	Artificial or Unknown found in <213> in SEQ ID (15)
W 213	Artificial or Unknown found in <213> in SEQ ID (16)
W 213	Artificial or Unknown found in <213> in SEQ ID (17)
W 213	Artificial or Unknown found in <213> in SEQ ID (18)
W 213	Artificial or Unknown found in <213> in SEQ ID (19)
W 213	Artificial or Unknown found in <213> in SEQ ID (20)

Input Set:

Output Set:

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Finished: 2007-09-26 16:48:27.466
Elapsed: 0 hr(s) 0 min(s) 12 sec(s) 724 ms
Total Warnings: 60
Total Errors: 0
No. of SeqIDs Defined: 92
Actual SeqID Count: 92

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (21)
W 213	Artificial or Unknown found in <213> in SEQ ID (22)
W 213	Artificial or Unknown found in <213> in SEQ ID (23)
W 213	Artificial or Unknown found in <213> in SEQ ID (24)
W 213	Artificial or Unknown found in <213> in SEQ ID (25)
W 213	Artificial or Unknown found in <213> in SEQ ID (26)
W 213	Artificial or Unknown found in <213> in SEQ ID (27)
W 213	Artificial or Unknown found in <213> in SEQ ID (28) This error has occurred more than 20 times, will not be displayed
W 402	Undefined organism found in <213> in SEQ ID (39)
W 402	Undefined organism found in <213> in SEQ ID (43)
W 402	Undefined organism found in <213> in SEQ ID (44)
W 402	Undefined organism found in <213> in SEQ ID (46)
W 402	Undefined organism found in <213> in SEQ ID (50)
W 402	Undefined organism found in <213> in SEQ ID (51)
W 402	Undefined organism found in <213> in SEQ ID (54)
W 402	Undefined organism found in <213> in SEQ ID (55)
W 402	Undefined organism found in <213> in SEQ ID (56)
W 402	Undefined organism found in <213> in SEQ ID (59)
W 402	Undefined organism found in <213> in SEQ ID (61)
W 402	Undefined organism found in <213> in SEQ ID (62) This error has occurred more than 20 times, will not be displayed

SEQUENCE LISTING

<110> Inouye, Masayori
 Zhang, Junjie
 Zhang, Yong Long
 Qing, Guoliang
 Suzuki, Motoo

<120> mRNA Interferases and Methods of Use Thereof

<130> University of Medicine & Dentistry of New Jersey (601-1-131PCT)

<140> 10560303

<141> 2007-09-26

<150> PCT/US2004/018571

<151> 2004-06-14

<150> 60/543,693

<151> 2004-02-11

<150> 60/478,515

<151> 2003-06-13

<160> 92

<170> FastSEQ for Windows Version 4.0

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<211> 336

<212> DNA

<213> E. coli

<400> 1

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aacaaaacag gtatgtgtct gtgtgttcct tgtacaacgc aatcaaaagg atatccgttc 180
gaagttgttt tatccggtca ggaacgtgat ggcgtagcgt tagctgatca ggtaaaaagt 240
atcgctggc gggcaagagg agcaacgaag aaaggaacag ttgccccaga ggaattacaa 300
ctcattaaag ccaaaattaa cgtactgatt gggtag 336
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<210> 2

<211> 111

<212> PRT

<213> E. coli

<400> 2

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Phe Asp Pro Thr Lys Gly Ser Glu Gln Ala Gly His Arg Pro Ala Val
              20              25              30
Val Leu Ser Pro Phe Met Tyr Asn Asn Lys Thr Gly Met Cys Leu Cys
              35              40              45
Val Pro Cys Thr Thr Gln Ser Lys Gly Tyr Pro Phe Glu Val Val Leu
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65	70	75
Ile Ala Trp Arg Ala Arg Gly Ala Thr Lys Lys Gly Thr Val Ala Pro		
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Glu Glu Leu Gln Leu Ile Lys Ala Lys Ile Asn Val Leu Ile Gly		
100	105	110

<210> 3
 <211> 333
 <212> DNA
 <213> E. coli

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 ggaacgcggc cgggtgctgat tgtcacaccg gcggccttta atcgcgtgac ccgcctgcct 120
 gttgttgtgc ccgtaaccag cggaggcaat tttgcccgca ctgccggctt tgcgggtgtcg 180
 ttggatggtg ttggcatacg taccacaggt gttgtacgtt gcgatcaacc ccggacaatt 240
 gatatgaaag cacggggcgg aaaacgactc gaacgggttc cggagactat catgaacgaa 300
 gttcttggcc gcctgtccac tattctgact tga 333

<210> 4
 <211> 110
 <212> PRT
 <213> E. coli

<400> 4
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 His Glu Gln Gln Gly Thr Arg Pro Val Leu Ile Val Thr Pro Ala Ala
 20 25 30
 Phe Asn Arg Val Thr Arg Leu Pro Val Val Val Pro Val Thr Ser Gly
 35 40 45
 Gly Asn Phe Ala Arg Thr Ala Gly Phe Ala Val Ser Leu Asp Gly Val
 50 55 60
 Gly Ile Arg Thr Thr Gly Val Val Arg Cys Asp Gln Pro Arg Thr Ile
 65 70 75 80
 Asp Met Lys Ala Arg Gly Gly Lys Arg Leu Glu Arg Val Pro Glu Thr
 85 90 95
 Ile Met Asn Glu Val Leu Gly Arg Leu Ser Thr Ile Leu Thr
 100 105 110

<210> 5
 <211> 249
 <212> DNA
 <213> E. coli

<400> 5
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 ttaatgcagg cgctcaatct gaattattgat gatgaagtga agattgacct ggtggatggc 120

aaattaatta ttgagccagt gcgtaaagag cccgtattta cgcttgctga actgggtcaac 180
gacatcacgc cggaaaacct ccacgagaat atcgactggg gagagccgaa agataaggaa 240
gtctggttaa 249

<210> 6
<211> 82
<212> PRT
<213> E. coli

<400> 6
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1 5 10 15
Ile Pro Ala Thr Leu Met Gln Ala Leu Asn Leu Asn Ile Asp Asp Glu
20 25 30
Val Lys Ile Asp Leu Val Asp Gly Lys Leu Ile Ile Glu Pro Val Arg
35 40 45
Lys Glu Pro Val Phe Thr Leu Ala Glu Leu Val Asn Asp Ile Thr Pro
50 55 60
Glu Asn Leu His Glu Asn Ile Asp Trp Gly Glu Pro Lys Asp Lys Glu
65 70 75 80
Val Trp

<210> 7
<211> 258
<212> DNA
<213> E. coli

<400> 7
atgcatacca cccgactgaa gaggggttggc ggctcagtta tgctgaccgt cccaccggca 60
ctgctgaatg cgctgtctct gggcacagat aatgaagttg gcatgggtcat tgataatggc 120
cggctgattg ttgagccgta cagacgcccg caatattcac tggctgagct actggcacag 180
tgtgatccga atgctgaaat atcagctgaa gaacgagaat ggctggatgc accggcgact 240
ggtcaggagg aaatctga 258

<210> 8
<211> 85
<212> PRT
<213> E. coli

<400> 8
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20 25 30
Val Gly Met Val Ile Asp Asn Gly Arg Leu Ile Val Glu Pro Tyr Arg
35 40 45
Arg Pro Gln Tyr Ser Leu Ala Glu Leu Leu Ala Gln Cys Asp Pro Asn
50 55 60
Ala Glu Ile Ser Ala Glu Glu Arg Glu Trp Leu Asp Ala Pro Ala Thr
65 70 75 80
Gly Gln Glu Glu Ile
85

<210> 9
 <211> 24
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> T54 to K77 fragment of E. coli MazE

<400> 9
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 Asn Ile Asp Trp Gly Glu Pro Lys
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<210> 10
 <211> 18
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> N60 to K77 fragment of E. coli MazE

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 1 5 10 15
 Pro Lys

<210> 11
 <211> 30
 <212> RNA
 <213> Artificial Sequence

<220>
 <223> synthetic RNA substrate

<400> 11
 uaagaaggag auauacauau gaaucaaauc 30

<210> 12
 <211> 50
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> single stranded oligonucleotide

<400> 12
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<210> 13
 <211> 50

<212> DNA
<213> Artificial Sequence

<220>
<223> single stranded oligonucleotide

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<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> synthetic oligonucleotide

<400> 14
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<210> 15
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> DNA primer

<400> 15
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<210> 16
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> DNA primer

<400> 16
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<210> 17
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> DNA primer

<400> 17
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<210> 18
<211> 24

<212> DNA
 <213> Artificial Sequence

 <220>
 <223> DNA primer

 <400> 18
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 <210> 19
 <211> 27
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> DNA primer

 <400> 19
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 <210> 20
 <211> 27
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> DNA primer

 <400> 20
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 <210> 21
 <211> 27
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> DNA primer

 <400> 21
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 <210> 22
 <211> 24
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> DNA primer

 <400> 22
 gatccccaca atgcggtgac gagt 24

 <210> 23
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> DNA primer

<400> 23
 cacgttggtcc actttgttca ccgc 24

<210> 24
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> DNA primer

<400> 24
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<210> 25
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> DNA primer

<400> 25
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<210> 26
 <211> 30
 <212> RNA
 <213> Artificial Sequence

<220>
 <223> antisense RNA

<400> 26
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<210> 27
 <211> 30
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> complementary DNA

<400> 27
 gatttgattc atatgtatat ctctttctta 30

<210> 28
 <211> 22
 <212> DNA
 <213> Artificial Sequence

<220>

<223> DNA primer	
<400> 28	
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<212> DNA	
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<220>	
<223> DNA fragment	
<400> 29	
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<210> 30	
<211> 15	
<212> DNA	
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<223> synthetic oligonucleotide	
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<213> Artificial Sequence	
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<400> 33
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<210> 34
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<212> DNA
<213> Artificial Sequence

<220>
<223> DNA primer

<400> 34
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<210> 35
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> DNA primer

<400> 35
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<210> 36
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> DNA primer

<400> 36
tcctctatgg agttactagt g 21

<210> 37
<211> 16
<212> DNA
<213> Artificial Sequence

<220>
<223> DNA primer

<400> 37
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<210> 38
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> DNA primer

<400> 38

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23

<210> 39

<211> 330

<212> DNA

<213> *Bacillus halodurans*

<400> 39

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ggTTTTgcg tggTTgtcc aattaccaga caacaaaag gttatccttt tgaaatagaa 180
ataccaccg ggttacctat tgaaggggtt attcttactg accaagtaa aagtctggat 240
tggagagcaa gaaactttca cattaagga caagcaccag aggaaactgt tactgattgt 300
ttacaactta ttcatacatt tttatcttaa 330

<210> 40

<211> 363

<212> DNA

<213> *Staphylococcus epidermidis*

<400> 40

atgattagaa gaggagatgt ttatttagcg gatttatac cagttcaagg gtctgaacaa 60
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gtaattgtag ctgcgattac tgatgggatt aataaagcga aaataccaac ccacgtagaa 180
attgaaaaga aaaagtataa attagacaaa gattcagtta ttcttcttga acaaattaga 240
acactagata aaaagcgttt aaaagaaaaa ttaacatttt tatcagagag taaaatgata 300
gaggttgata atgccttaga tattagtttg ggattaaata actttgatca tcataaatct 360
taa 363

<210> 41

<211> 411

<212> DNA

<213> *Staphylococcus aureus*

<400> 41

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gttattgttg cggcaataac tggtaggatt aataaagcga aaataccgac acatgtagag 180
attgaaaaga aaaagtataa gttggataaa gactcagtta tattattaga acaaattcgt 240
acacttgata aaaaacgatt gaaagaaaaa ctgacgtact tatccgatga taaaatgaaa 300
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<210> 42

<211> 351

<212> DNA

<213> *Bacillus subtilis*

<400> 42

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actgctattg ttgcagccat aacagcacia atacagaaag cgaaattacc aaccacgtc 180
gaaatcgatg caaaacgcta cggTTTTgaa agagattccg ttattttgct ggagcaaatt 240
cggacgattg acaagcaaag gttaacggat aagattactc atctggatga tgaaatgatg 300
gataaggttg atgaagcctt acaaatcagt ttggcactca ttgattTTa g 351

<210> 43
<211> 324
<212> DNA
<213> *Neisseria meningitides*

<400> 43
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gaaatcaaaa agacacgtcc ttgtgtcgta gtctctcctc ctgaaataca caactatctc 120
aagactgtgc tgatcgttcc catgacgagc ggaagccgtc ctgccccgtt ccgcgtcaat 180
gtccgcctttc aggataaaga cggtttgctt ttgcccgaac agattagggc tgtggataaa 240
gccggattgg tcaaacatct tggcaattta gacaacagta cggctgaaaa actgtttgca 300
gtattgcagg agatgtttgc ctga 324

<210> 44
<211> 366
<212> DNA
<213> *Morganella morganii*

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ccggctgctt ttaaccgcgt gaccgcctg cctgttggtg tgcccgtagc cagcggaggt 180
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<210> 45
<211> 321
<212> DNA
<213> *Mycobacterium tuberculosis*

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caacgcgggc ggcgctacgc cgtggctcctc agccccggct cgatgccgtg gagtgtagta 120
accgtggtgc cgacgtcgac aagcgcccaa cctgcgggtt tccgaccaga gctggaagtc 180
atgggaacaa agacacgggt cctggtggat cagatccgga cgatcggcat cgtctatgtg 240
cacggcgatc cggtcgacta tctggaccgt gaccaaattg ccaaggtgga acacgccgtg 300
gcacgatacc ttggtctgtg a 321

<210> 46
<211> 109
<212> PRT
<213> *Bacillus halodurans*

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Pro Lys Leu Phe Asn Lys Asn Thr Gly Phe Ala Val Val Cys Pro Ile
35 40 45
Thr Arg Gln Gln Lys Gly Tyr Pro Phe Glu Ile Glu Ile Pro Pro Gly
50 55 60
Leu Pro Ile Glu Gly Val Ile Leu Thr Asp Gln Val Lys Ser Leu Asp
65 70 75 80

